Seq. I.D. No. 1

	90	100	109	118		128	138	148	
	92	102	112	122		132	142	152	
	89	99	109	119	CHO	129	139	149	
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Clq a	QPRPA	FSAIRRNF	PMGGNVVIPD	TVITNĢEE	PYQNH	SGRFV	CTVPGYYYFTF	QVLSQWEICLS	ivsss
Clq b	TQKIA	FSATRTINVE	LRRDQTIRFD	HVITNMNN	NYEPR	SGKFT	CKVPGLYYFTY	HASSRGNLCVN	LMRG-
Clq c	KFOSV	FTVTROTHOR	PAPNSLIRFN	AVLTNPQG	DYDTS	TGKFT	CKVPGLYYFVYI	HASHTANLCVL	LYRS-
	159	168	178	188		198	208	218	
	162	171	180	190		200	210	219	
	158	165	174	184		194	204	213	
	1		1	1			ŀ	1	
Clq a	RGQVR	RSLGFCDTTN	KGLFQVVSGG	MVLQLQQG	DQVWV	EKDPK	KGHIYQGSEAD	SVFSGFLIFPS	A
Clq b	RERAQ	KVVTFCDYAY	N-TFQVTTGG	MVLKLEQG	ENVFL	QATDK	NSLLGMEG-AN	SIFSGFLLFPD	MEA
Clq_c	-GV	KVVTFCGHTS	K-TNQVNSGG	VLLRLQVG	EEVWL	AVNDY	YDMVGIQG-SD	SVFSGFLLFPD	

Amino acid sequence of gClq domains aligned with structural similarity.

Seq. I.D. No. 2

Amino acid sequence of ClqA

Complement component 1, q subcomponent, alpha polypeptide precursor ACCESSION AAH30153 GI:20988805

1 MEGPRGWLVL CVLAISLASM VTEDLCRAPD GKKGEAGRPG RRGRPGLKGE QGEPGAPGIR
61 TGIQGLKGDQ GEPGPSGNPG KVGYPGPSGP LGARGIPGIK GTKGSPGNIK DQPRPAFSAI
121 RRNPPMGGNV VIFDTVITNQ EEPYQNHSGR FVCTVPGYYY FTFQVLSQWE ICLSIVSSSR
181 GQVRRSLGFC DTTNKGLFQV VSGGMVLQLQ QGDQVWVEKD PKKGHIYQGS EADSVFSGFL
241 IFPSA

Seq. I.D. No. 3

Amino acid sequence of C1qB

complement component 1, q subcomponent, beta polypeptide precursor ACCESSION NP_000482 GI:11038662

- 1 MMMKIPWGSI PVLMLLLLG LIDISQAQLS CTGPPAIPGI PGIPGTPGPD GQPGTPGIKG 61 EKGLPGLAGD HGEFGEKGDP GIPGNPGKVG PKGPMGPKGG PGAPGAPGPK GESGDYKATQ 121 KIAFSATRTI NVPLRRDQTI RFDHVITNMN NNYEPRSGKF TCKVPGLYYF TYHASSRGNL 181 CVNLMRGRER AQKVVTFCDY AYNTFQVTTG GMVLKLEQGE NVFLQATDKN SLLGMEGANS
- 241 IFSGFLLFPD MEA

Seq. I.D. No. 4

Amino acid sequence of C1qC

Complement Clq subcomponent, C chain precursor. ACCESSION P02747 GI:20178281

1 MMMKIPWGSI PVLILLLLG LIDISQAQLS CTGPPAIPGI PGIPGTPGPD GQPGTPGIKG
61 EKGLPGLAGD HGEFGEKGDP GIPGNPGKVG PKGPMGPKGG PGAPGAPGPK GESGDYKATQ
121 KIAFSATRTI NVPLRRDQTI RFDHVITNMN NNYEPRSGKF TCKVPGLYYF TYHASSRGNL
181 CVNLMRGRER AQKVVTFCDY AYNTFQVTTG GMVLKLEQGE NVFLQATDKN SLLGMEGANS
241 IFSGFLLFPD MEA